



Figure 2-1

AccI		GT'mk_AC (0 Err) - 2 Fragment(s)
	243	909
Apal		G_GGCC'C (0 Err) - 2 Fragment(s)
	402	750
BamHI		G'GATC_C (0 Err) - 2 Fragment(s)
	200	952
EcoRI		G'AATT_C (0 Err) - 2 Fragment(s)
	343	809
HindIII		A'AGCT_T (0 Err) - 2 Fragment(s)
	135	1017
NciI		CC's_GG (0 Err) - 2 Fragment(s)
	196	956
NcoI		C'CATG_G (0 Err) - 4 Fragment(s)
	60	135 288 669
PstI		C_TGCA'G (0 Err) - 3 Fragment(s)
	337	366 449
PvuII		CAG'CTG (0 Err) - 2 Fragment(s)
	570	582
Sall		G'TCGA_C (0 Err) - 2 Fragment(s)
	242	910

Linear Map of Sequence:

1 gggcaggcaatgagagctgcactctggctgggaaggcatgagtgacacagaccacagca
 ccccgccgttactctcgacgtgagaccgacccttcgtactcactgtctgggtgcgt * 60
 * * * * * * * * * * * *
 61 aggccgtgggtaagtgtggacctttgttaccagagagaacatcatggtggcttcaa
 tccggccacccattcacacacctggaaacacatggtctctttgttagtaccaccgaaat * 120
 * * * * * * * * * * * *
 HindIII
 \
 21 ggggtctggactcaagcttctggaaagcagtcacagcggaaatttctggccatgcttatt
 ccccagacctgagttcgaaaagaccttcgtcagtgtcgccctaaagaccggtaacgaaataa * 180
 * * * * * * * * * * * *
 BamHI
 \
 181 ttgttctccctcagcctggatccaccatcaactggggtggaacagaaaaaggcttaccg
 aaacaagaggagtcggacccttaggttgttagttgacccaccccttgccttcggaaatggc * 240
 * * * * * * * * * * * *
 SalI

Figure 2-2

Figure 2-3

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Citation

Algorithm Citation:

Not given.

Program Citation:

Program Citation:
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